

Figure 1a

6A1 VH			
6A1	EVQLVQSGAEVKKPGESLTISCKGSGYNFE	NKWTIG	WVRQMPGKGLEWVG
Vh5_DP-73__5-51	EVQLVQSGAEVKKPGESLKISCKGSGYSFT	SNWTIG	WVRQMPGKGLEWVG
6A1	LYPGRDSIDTRVSPSFOG		
Vh5_DP-73__5-51	QVTISVDKISISTAYLQWSSLKASDTAMYICAR	SH	100
6A1	WGRGTMVTVSS	128	SEQ ID NO:108
Vh5_DP-73__5-51	WGQGTMTVTVSS	JH3	SEQ ID NO:144
6A1 VL			
6A1	SSELTQDPAVSVALGQTVRITC	OGDSIERSVAVAS	WYQQKPGQAPVLIY
Vlambda3_DPL16__31_	SSELTQDPAVSVALGQTVRITC	OGDSIERSVAVAS	WYQQKPGQAPVLIY
6A1	NKRIPS	GIPDRFSGSSSGNTASLTITGAQAEDEADYYC	HSRDSGSHVVL
Vlambda3_DPL16__31_	NNRIPS	GIPDRFSGSSSGNTASLTITGAQAEDEADYYC	NSRDSGSHVVL
6A1	GGTKLTVLGA	110	SEQ ID NO:109
Vlambda3_DPL16__31_	GGTKLTVL..	JL3	SEQ ID NO:151

Figure 1b

7A2 VH

7A2
Vh5_DP-73__5-51_
GVQLVQSGAEVKKPESLTISKSGYNFF NAWTC WVRQMPGKGLEWVG 50
EVQLVQSGAEVKKPESLTISKSGYSFT SMTTC WVRQMPGKGLEWVG 50
7A2
Vh5_DP-73__5-51_
EYPTDSDDTRVSPSFQGV BVTISYDKSISTAYLQWSSLKASDTAMYCAR STJ 100
INPCDSDTRVSPSFQGV QVTISADKSISTAYLQWSSLKASDTAMYCAR ..
7A2
Vh5_DP-73__5-51_
RYCPGGGROMSGIACMDV WGQGTMTVSS 128 SEQ ID NO:110
..... WGQGTMTVSS JH3 SEQ ID NO:144

7A2 VL

7A2
Vlambda3_DPL16__31_
SSELTQDPAVSVALGQTVRITC QGDSIRSWAVN WFQQKPGQAPLLVYV AK 50
SSELTQDPAVSVALGQTVRITC QGDSIRSWAVS WYQQKPGQAPLVLYY GK 50
7A2
Vlambda3_DPL16__31_
NKRPS GIPDRFSGSSSGNTASLTITGAQAEDADYYC NSRDSSGNEHVA FG 100
NNRPS GIPDRFSGSSSGNTASLTITGAQAEDADYYC NSRDSSGNEHVA FG 100
7A2
Vlambda3_DPL16__31_
GGTKLTVLGA 110 SEQ ID NO:111
GGTKLTVL.. JL3 SEQ ID NO:151

Figure 1c

7A4 VH					
7A4					
Vh5_DP-73__5-51_					
	EVQLVQSGAEVKKPGESLTISCKGSGYNFF	NVWITG	WVRQMPGKDLWMG	50	
	EVQLVQSGAEVKKPGESLKISCKGSGYSFT	SRWITG	WVRQMPGKGLEWMG	50	
7A4					
Vh5_DP-73__5-51_					
	IVPTDSDFRYSPSPFOG	QVTISVDKSI	TAYLQWSSLKASDTAMYYCAR	100	
	IVPGDSDFRYSPSPFOG	QVTISADKSI	TAYLQWSSLKASDTAMYYCAR	100	
7A4					
Vh5_DP-73__5-51_					
	RVGPGGRGYSQWGMDDV	WGQGTMTVSS	128	SEQ ID NO:112	
	WGQGTMTVSS	JH3	SEQ ID NO:144	
7A4 V					
7A4					
Vlambda3_DPL16__31_					
	SSELTQDPAVSVALGQTVRITC	RGDSIRNNWAS	WYQQKPGQAPVLVIY	50	
	SSELTQDPAVSVALGQTVRITC	OGDSIRSNWAS	WYQQKPGQAPVLVIY	50	
7A4					
Vlambda3_DPL16__31_					
	NNRBS	GIPDRFSGSSSGNTASLTITGAQAEDADYYC	NSRDSSGNHMY	FG 100	
	NNRBS	GIPDRFSGSSSGNTASLTITGAQAEDADYYC	NSRDSSGNHMY	FG 100	
7A4					
Vlambda3_DPL16__31_					
	GGTKLTVLGA	110	SEQ ID NO:113		
	GGTKLTVL..	JL3	SEQ ID NO:151		

Figure 1d

7A5 VH			
7A5			
Vh5_DP-73__5-51_	GVQLVESGAEVKKPGESLTISKSGYNFF	NMMITG	WVRQMPGKGLEWMG II 50
	EVQLVQSGAEVKKPGESLKISKSGYSFT	SMWITG	WVRQMPGKGLEWMG II
7A5			
Vh5_DP-73__5-51_	IMPTDSDTRMSPSFOG	QVTISVDKISISTAYLQWSSLKASDTAMYICAR	SI 100
	IMPGDSDDTRMSPSFOG	QVTISADKISISTAYLQWSSLKASDTAMYICAR	..
7A5			
Vh5_DP-73__5-51_	RACPGGRCVSGIAGMDV	WGRGTLVTVSS 128	SEQ ID NO:114
	WGRGTLVTVSS JH2	SEQ ID NO:145
7A5 VL			
7A5			
Vlambda3_DPL16__31_	SSELTQDPAVSVALGQTVRITC	QGDSTRSMVAS	WYQQKPGQAPVLVIY CK 50
	SSELTQDPAVSVALGQTVRITC	QGDSTRSMVAS	WYQQKPGQAPVLVIY CK
7A5			
Vlambda3_DPL16__31_	NNRPS	GIPDRFSGSSSGNTASLTITGAQAEDEADYYC	NSRDSGNGHW FG 100
	NNRPS	GIPDRFSGSSSGNTASLTITGAQAEDEADYYC	NSRDSGNGHW FG
7A5			
Vlambda3_DPL16__31_	GGTKLTVLGA 110	SEQ ID NO:115	
	GGTKLTVL.. JL3	SEQ ID NO:151	

Figure 1e

7A6 VH

7A6	EVQLVQSGAEVKKPGESLTISKSGSYNFE	NVMTIC	WVRQMPGKGLEWMG	50
Vh5_DP-73__5-51_	EVQLVQSGAEVKKPGESLKISKSGSYFT	SYMTIC	WVRQMPGKGLEWMG	
7A6	QVTISYDKSISTAYLQWSSSLKASDTAMYCAR	ST		100
Vh5_DP-73__5-51_	QVTISADKSISTAYLQWSSSLKASDTAMYCAR	..		
7A6	WGQGTTLVTSS	128	SEQ ID NO:116	
Vh5_DP-73__5-51_	WGQGTTLVTSS	JH5	SEQ ID NO:146	

7A6 VL

7A6	SSELTQDPAVSVALGQTVRITC	QGDSTRSNTIN	WEQQKPGQAPLLVY	50
Vlambda3_DPL16__31_	SSELTQDPAVSVALGQTVRITC	QGDSTRSNTAS	WEQQKPGQAPLVII	
7A6	GIPDRFSGSSSGNTASLTITGAQAEDEADYYC	NSRDSSGNEHVA	FG	100
Vlambda3_DPL16__31_	GIPDRFSGSSSGNTASLTITGAQAEDEADYYC	NSRDSSGNEHVA	FG	
7A6	GGTKLTVL	108	SEQ ID NO:117	
Vlambda3_DPL16__31_	GGTKLTVL	JL3	SEQ ID NO:151	

Figure 1f

8A1 VH				
8A1				
Vh5_DP-73__5-51_		EVQLVQSGAEVKKPESLTISKCKPGYNFF	NMMWIG	WVRQMPGKGLEWMG 50
		EVQLVQSGAEVKKPESLKISKCKSGYSFT	SMWIG	WVRQMPGKGLEWMG 50
8A1				
Vh5_DP-73__5-51_		QVTISYDKSISTAYLQWSSLKASDTAMYICAR	SM	100
		QVTISADKSISTAYLQWSSLKASDTAMYICAR	..	
8A1				
Vh5_DP-73__5-51_		WGQGTMTVTSS 128	SEQ ID NO:118	
		WGQGTMTVTSS JH3	SEQ ID NO:144	
8A1 VL				
8A1				
Vlambda3_DPL16__31_		SSELTQDPAVSVALGQTVRITC	QCDLSRSMVAS	WYQQKPGQAPVLVIY 50
		SSELTQDPAVSVALGQTVRITC	QCDLSRSMVAS	WYQQKPGQAPVLVIY 50
8A1				
Vlambda3_DPL16__31_		GIPDRFSGSSSGNTASLTITGAQAEDEADYYC	NSRDSSGNEHW	FG 100
		GIPDRFSGSSSGNTASLTITGAQAEDEADYYC	NSRDSSGNEHW	FG 100
8A1				
Vlambda3_DPL16__31_		GGTKLTVLGA 110	SEQ ID NO:119	
		GGTKLTVL.. JL3	SEQ ID NO:151	

Figure 1g

9A2 VH			
9A2	QVQLVQSGAEVRKPGASVKVCKTSGYTR	NVDIN	WVRQAPQGQLEWVG R 50
Vh1_DP-14__1-18_	QVQLVQSGAEVKKPGASVKVCKASGYTFT	SYGLIS	WVRQAPQGQLEWVG W
9A2	TSCHYGNTDTHAOKFOG	RETMTKDTSTSTAYMELRSLTFDDTA	VYYCAR SQ 100
Vh1_DP-14__1-18_	TSAYVNGNTNVAOKLQG	RVTMTTDTSTSTAYMELRSLRSDDTA	VYYCAR ..
9A2	WNVDM	WGRGTLVTVSS 116	SEQ ID NO:120
Vh1_DP-14__1-18_	WGRGTLVTVSS JH2	SEQ ID NO:147
9A2 VL			
9A2	NFMLTQPHSVSESPGKTVTISC	TRSSGSAISNVAQ	WYQORPGSSPTTVIF 50
Vlambda6_6a	NFMLTQPHSVSESPGKTVTISC	TRSSGSAISNVAQ	WYQORPGSSPTTVIY
9A2	EDNRRPS	GVPDRFSGSIDTSSNSASLTISGLKTE	DEADYYC QSPDSTNLY 100
Vlambda6_6a	EDNORPS	GVPDRFSGSIDSSNSASLTISGLKTE	DEADYYC QSPDSSNLY
9A2	V	FCGGTKLTVLGA 113	SEQ ID NO:121
Vlambda6_6a	V	FCGGTKLTVL.. JL2/JL3	SEQ ID NO:152

Figure 1h

11A1 VH			
11A1	EVQLVESGGGVVQPGRSLRLSCAASGFTFS	DEAME	WVRQIPGKGLEWLS
Vh3_DP-50__3-33_	QVQLVESGGGVVQPGRSLRLSCAASGFTFS	SMGME	WVRQAPGKGLEWVA
11A1	LRHDCSTAVAVAGSVKGC		
Vh3_DP-50__3-33_	LNWIDGSNNKAVADSVMKGC		
11A1	CSGCPHAFEPV		
Vh3_DP-50__3-33_	WGKGLTVTVSS	121	SEQ ID NO:122
	WGQGLTVTVSS	JH1/JH4/JH5
			SEQ ID NO:148
	* Vernier residue		
11A1 VL			
11A1	SYVLTQPPSASGTPGQRTISC	SGSNSNIGCTATVN	WFQQLPGTAPKLLIY
Vlambda1_DPL2__1c_	QSVLTQPPSASGTPGQRTISC	SGSSSNIGCSNFTVN	WYQQLPGTAPKLLIY
11A1	SNNORPS		
Vlambda1_DPL2__1c_	SNNORPS		
11A1	FGGGTKVTVLGA	112	SEQ ID NO:123
Vlambda1_DPL2__1c_	FGIGTKVTVL	JL1	SEQ ID NO:154

Figure 1i

11A2 VH				
11A2	EVQLLESGGGLVQPGGSLRLS	CAASGFTFS	SNVAMS	WVRQAPGKGLEWVS
Vh3_DP-47_3-23_	EVQLLESGGGLVQPGGSLRLS	CAASGFTFS	SNVAMS	WVRQAPGKGLEWVS
11A2	TSVSGSGSTVWADSVKGC			
Vh3_DP-47_3-23_	TSVSGSGSTVWADSVKGC			
11A2	WGQGTMTVSS	127	SEQ ID NO:124	
Vh3_DP-47_3-23_	WGQGTMTVSS	JH3	SEQ ID NO:149	
11A2 VL				
11A2	SSELTQDPDVSMALGQTVTISC			
Vlambda3_DPL16_31_	SSELTQDPDVSMALGQTVTISC			
11A2	GIPDRFSGSDSGDTASLTITGAQAEDEGDIYC			
Vlambda3_DPL16_31_	GIPDRFSGSDSGDTASLTITGAQAEDEGDIYC			
11A2	SGTKVTVLGA	110	SEQ ID NO:125	
Vlambda3_DPL16_31_	TGTKVTVL..	JL1	SEQ ID NO:153	

Figure 1j

11A3 VH

Query
Vh1_DP-14_1-18_ EVQLVQSGAEVKKPGASVKVSCKASGYST NYGLN WVRQAPGGLEWMG W 50
QVQLVQSGAEVKKPGASVKVSCKASGYTFT SMGLS WVRQAPGGLEWMG W

Query
Vh1_DP-14_1-18_ ISPYTGTGNTNNAOKFOG RVTMTTDKSTSTAYMDLRSLRSDDTAVYYCAR ERI 100
ISAVNCGNTNNAOKILOG RVTMTTDISTSTAYMELRSLRSDDTAVYYCAR ..

Query
Vh1_DP-14_1-18_ ESHCITGGSCMPEDS WGRGTLTVSS 125 SEQ ID NO:126
..... WGRGTLTVSS JH2 SEQ ID NO:147

11A3 VL

Query
Vlambda3_DPL16_31_ SSELTDQDPAVSVALGQTVRITC QCDSEIRNNVAS WYQQKPGQAPLLVMF GK 50
SSELTDQDPAVSVALGQTVRITC QCDSEIRSVVAS WYQQKPGQAPVLVIY GK

Query
Vlambda3_DPL16_31_ NNRPS EIPGRFSGSSSGNTASLTITGAQAEEADYYC NSRDRNSHOWV FG 100
NNRPS GIPDRFSGSSSGNTASLTITGAQAEEADYYC NSRDSGNEHW FG

Query
Vlambda3_DPL16_31_ GGKLTVLGA 110 SEQ ID NO:127
GGKLTVL.. JL3 SEQ ID NO:151

Figure 1k

```

11A4 VH
11A4
Vh3_DP-47__3-23_ EVQLLESGGGLVQPGGSLRLSCAASGFTFS SWAMS WVRQAPGKGLEWVS A 50
EVQLLESGGGLVQPGGSLRLSCAASGFTFS SWAMS WVRQAPGKGLEWVS A
11A4
Vh3_DP-47__3-23_ TSGSGGSGTAYADSVKQ RFTISRDN SKNTLYLQMN SLRAEDTAVYYC SP 100
TSGSGGSGTAYADSVKQ RFTISRDN SKNTLYLQMN SLRAEDTAVYYCAK ..
11A4
Vh3_DP-47__3-23_ YSGRWYSEDP WGQGTMTVTVSS 121 SEQ ID NO:128
YSGRWYSEDP WGQGTMTVTVSS JH3 SEQ ID NO:149
..... WGQGTMTVTVSS JH3
* Vernier residue

11A4 VL
11A4
Vlambda3_DPL23__3r_ SYELTQPPSVSVSPGQTATITC SGDDPLCNKAMS WYQQKPGQSPVLVIY OD 50
SYELTQPPSVSVSPGQTASITC SGDKTGDKNVAC WYQQKPGQSPVLVIY OD
11A4
Vlambda3_DPL23__3r_ TKRPS GIPERFSGNSGNIATLTISGTQAVDEADYYC QVMDTGHVW FGCG 100
SKRPS GIPERFSGNSGNTATLTISGTQAMDEADYYC QAWDSSTA FGCG
11A4
Vlambda3_DPL23__3r_ TKLTVLGA 108 SEQ ID NO:129
TKLTVL.. JL3 SEQ ID NO:155

```

Figure 11

11A5 VH					
Query					
Vh1_DP-14__1-18__					50
	QVQLVQSGAEVKKPGASVKVSCKASGYST	NYGYN	WVRQAPGQGLEWMG		
	QVQLVQSGAEVKKPGASVKVSCKASGYTFT	SYGTS	WVRQAPGQGLEWMG		
Query					100
Vh1_DP-14__1-18__					
	RVMTMTDKSTSTAYMDLRLSLRSDDDTAVYYCAR	ET			
	RVMTMTDTSTSTAYMELRLSLRSDDDTAVYYCAR				
Query					
Vh1_DP-14__1-18__					
	WGKGLTVTVSS	125	SEQ ID NO:130		
	WGQGLTVTVSS	JH1	SEQ ID NO:145		
11A5 VL					
Query					50
Vlambda3_DPL16__31__					
	SSELTQDPAVSVALGQTVRITC	QCDSLRSVAAS	WYQKPGQAPVLVIY	CK	
	SSELTQDPAVSVALGQTVRITC	QCDSLRSVAAS	WYQKPGQAPVLVIY	CK	
Query					100
Vlambda3_DPL16__31__					
	GIPDRFSGSSSGNTASLTITGAQAEADYYC	NSRDSGNNHHMV		F	
	GIPDRFSGSSSGNTASLTITGAQAEADYYC	NSRDSGNNH		F	
Query					
Vlambda3_DPL16__31__					
	GGGKTVTVLGA	111	SEQ ID NO:131		
	GGGKLTVL..	JL3	SEQ ID NO:151		

Figure 1m

11A11 VH		
11A11	QVQLVESGGGLVKPGGSLRLS	CAASGFTFS SHTMN WVRQAQCKGLEWVS 50
Vh3_DP-77_3-21_	EVQLVESGGGLVKPGGSLRLS	CAASGFTFS SYSMN WVRQAQCKGLEWVS 50
11A11	TSGSGRTAVASDSVKG	RFTISRDAAKNSLYLQMN
Vh3_DP-77_3-21_	TSSSSSMDVADSVKG	RFTISRDNAKNSLYLQMN
11A11	FGDMLFDS	WGQGLTVTVSS 119 SEQ ID NO:132
Vh3_DP-77_3-21_	WGQGLTVTVSS JH1/JH4/JH5 SEQ ID NO:150
11A11 VL		
11A11	NFMLTQPHSVSQSPGKTVTISC	TPSSGRTFASNEVQ WYQORPGSAPTIVY 50
Vlambda6_6a	NFMLTQPHSVSES	PGKTVTISC PRSSGRTFASNNVQ WYQORPGSPTTVIY
11A11	EDNRPRPS	GVPDRFSGSIDSSSNSASLTISGLKTEDEADYYC OSMDARNQV 100
Vlambda6_6a	EDNORPS	GVPDRFSGSIDSSSNSASLTISGLKTEDEADYYC OSMDSSN 100
11A11	FGTGTKVTVLGA 112	SEQ ID NO:133
Vlambda6_6a	FGTGTKVTVL.. JL1	SEQ ID NO:156

[illegible]

[illegible]

Figure 1p

11A7 VH			
11A7			
Vh1_DP-14__1-18__	EVQLVQSGAEVKKPGASVKVSCKASGYST	NYGLD	WVRQAPQGLEWVG 50
	QVQLVQSGAEVKKPGASVKVSCKASGYTFT	SWGTS	WVRQAPQGLEWVG
11A7			
Vh1_DP-14__1-18__	TSPTTCMTNVAQKFOG		
	TSANGNFNVAQKFOG		
	RVTMTTDKSTSTAYMDLRLSLRSDDTAVYYCAR	PH	100
	RVTMTTDISTSTAYMELRLSLRSDDTAVYYCAR		..
11A7			
Vh1_DP-14__1-18__	FSHCIGGSCMPFDS		
	WGRGTMVTVSS 125	SEQ ID NO:136	
 WGRGTLVTVSS JH2	SEQ ID NO:147	
11A7 VL			
11A7			
Vlambda3_DPL16__31__	SSELTQDPAVSVALGQTVRITC	QGDSESRNVAAS	WYQQKPGQAPVLVIY GK 50
	SSELTQDPAVSVALGQTVRITC	QGDSESRNVAAS	WYQQKPGQAPVLVIY GK
11A7			
Vlambda3_DPL16__31__	NNRPS	GIPDRFSGSSSGNTASLTITGAQAEDEADYYC	NSRDSGSGNHNWV 100
	NNRPS	GIPDRFSGSSSGNTASLTITGAQAEDEADYYC	NSRDSGSGNEL 100
11A7			
Vlambda3_DPL16__31__	FGGGTKVTVLGA 112	SEQ ID NO:137	
	FGGGTKLTVL.. JL3	SEQ ID NO:151	

[illegible]

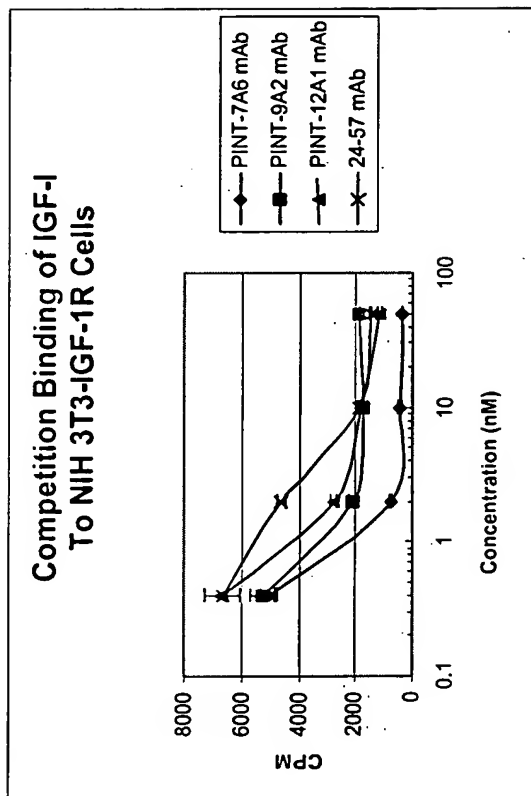
Figure 1r

12A3 VH			
12A3			
Vh1_DP-14_1-18_	QVQLVQSGAEVKKPGASVKVSKKASGYST	NYGLEN	WVRQAPGQGLEWMG W 50
	QVQLVQSGAEVKKPGASVKVSKKASGYTFT	SYGHS	WVRQAPGQGLEWMG W
12A3			
Vh1_DP-14_1-18_	TSPMTGNTNVAOKFOG	RVTMTSDKSTSTAYMDLRLSLRSDDTAIYYCAR	ETI 100
	TSVANGNTNVAOKFOG	RVTMTTDTSTSTAYMELRLSLRSDDTAVYYCAR ..	
12A3			
Vh1_DP-14_1-18_	FSHGSGCSGYPEDY	WGQGLTVTVSS 125	SEQ ID NO:140
	WGQGLTVTVSS JH1	SEQ ID NO:147
12A3 VL			
12A3			
Vlambda3_DPL16_31_	SSELTQDPAVSVALGQTVRITC	QGDSTRSWAS	WYQKPGQAPLVLIY GR 50
	SSELTQDPAVSVALGQTVRITC	QGDSTRSWAS	WYQKPGQAPLVLIY GK
12A3			
Vlambda3_DPL16_31_	NNRPS	GIPDRFSGSSSGNTASLTITGAQAEDADYYC	NSRDSSTNHGNWV 100
	NNRPS	GIPDRFSGSSSGNTASLTITGAQAEDADYYC	NSRDSSTNHGNWV
12A3			
Vlambda3_DPL16_31_	FGGGTQLTVLSA 112	SEQ ID NO:141	
	FGGGTQLTVL.. JL7	SEQ ID NO:157	

[illegible]

Figure 2

a



b

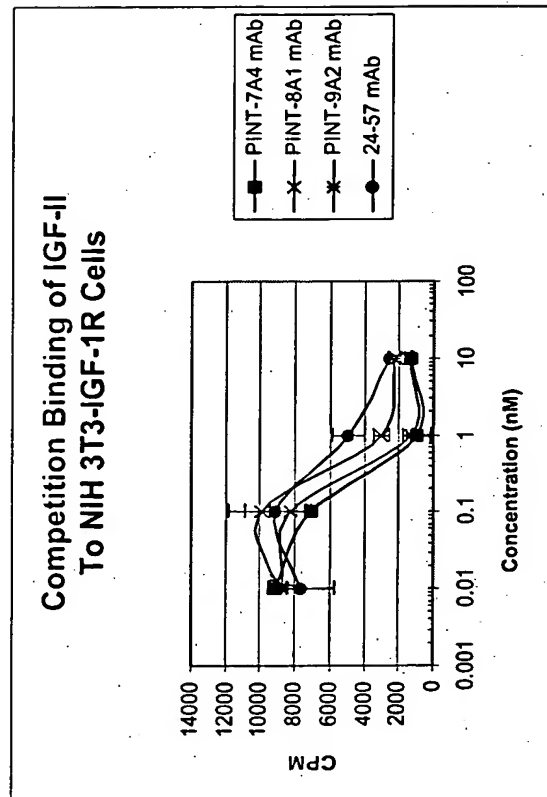


Figure 3

IGF-1R mAbs effect on insulin / receptor binding

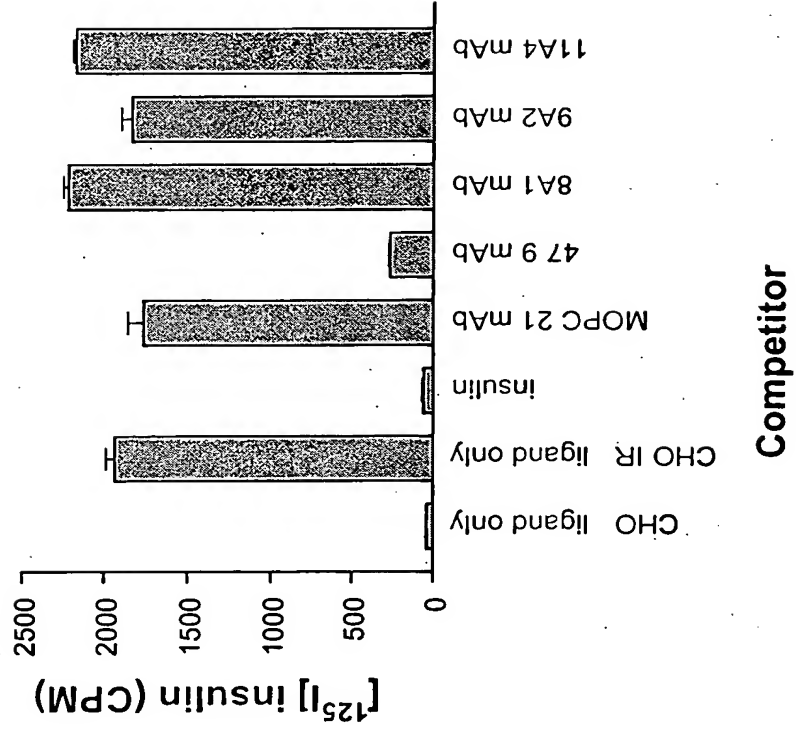


Figure 4

IGF-1R mAbs effect on insulin receptor activation

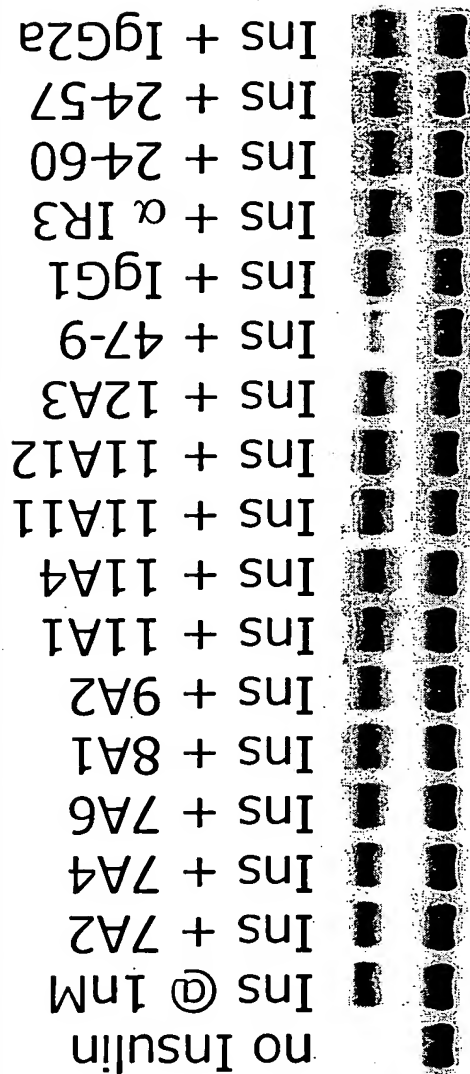


Figure 5

Saturable and specific binding
of IGF-1R mAbs - 3T3 hu-IGF1R
fibroblasts

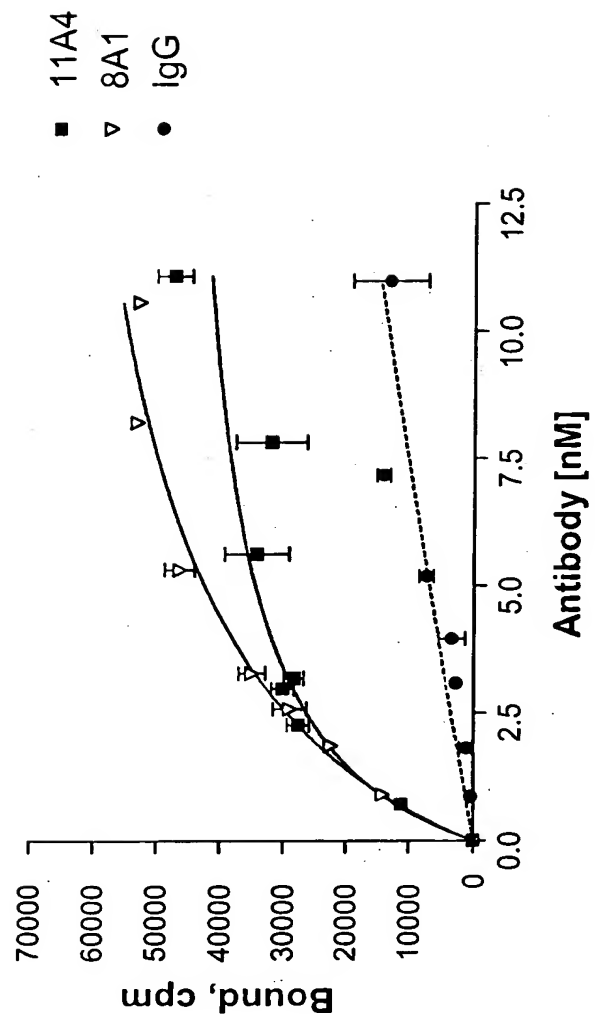


Figure 6

IGF-1R mAbs block cell proliferation - BrdU

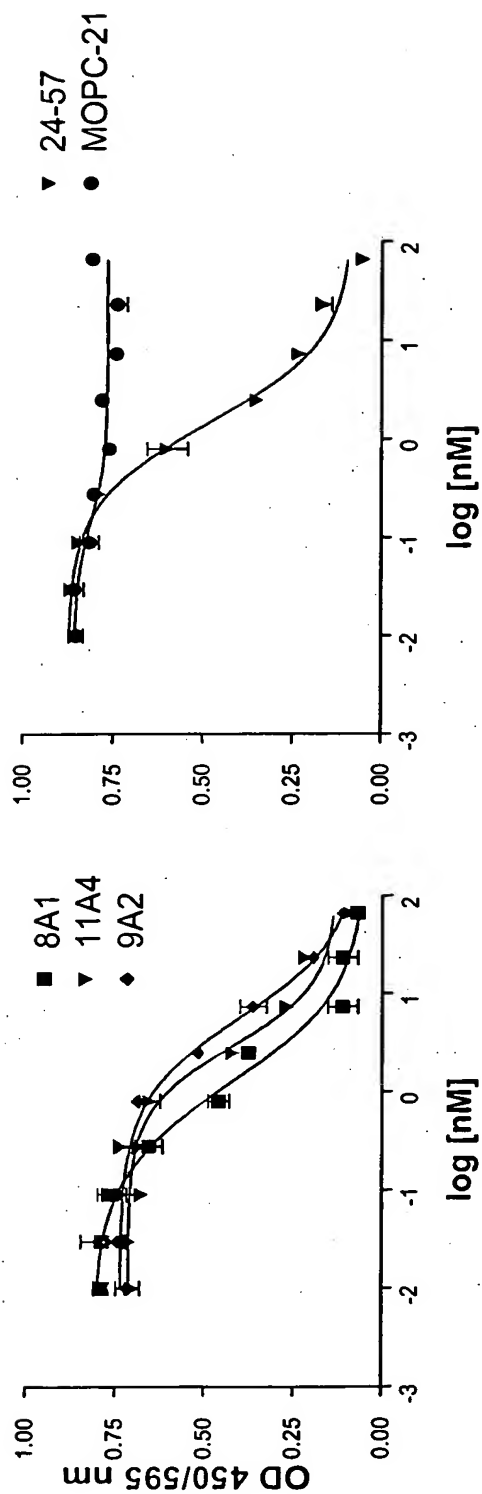


Figure 7

IGF-1R mAbs – receptor phosphorylation – Western blotting

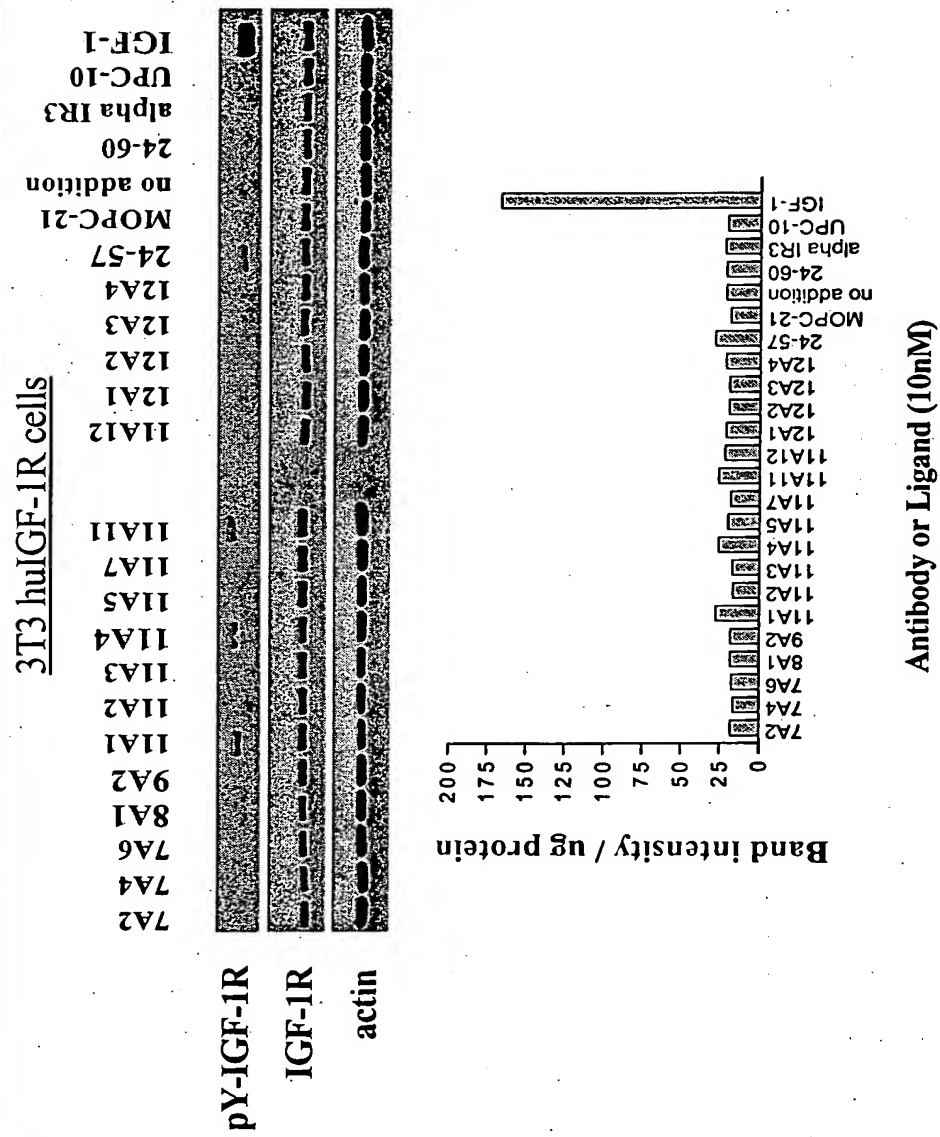


Figure 8

IGF-1R mAbs – receptor phosphorylation – ELISA

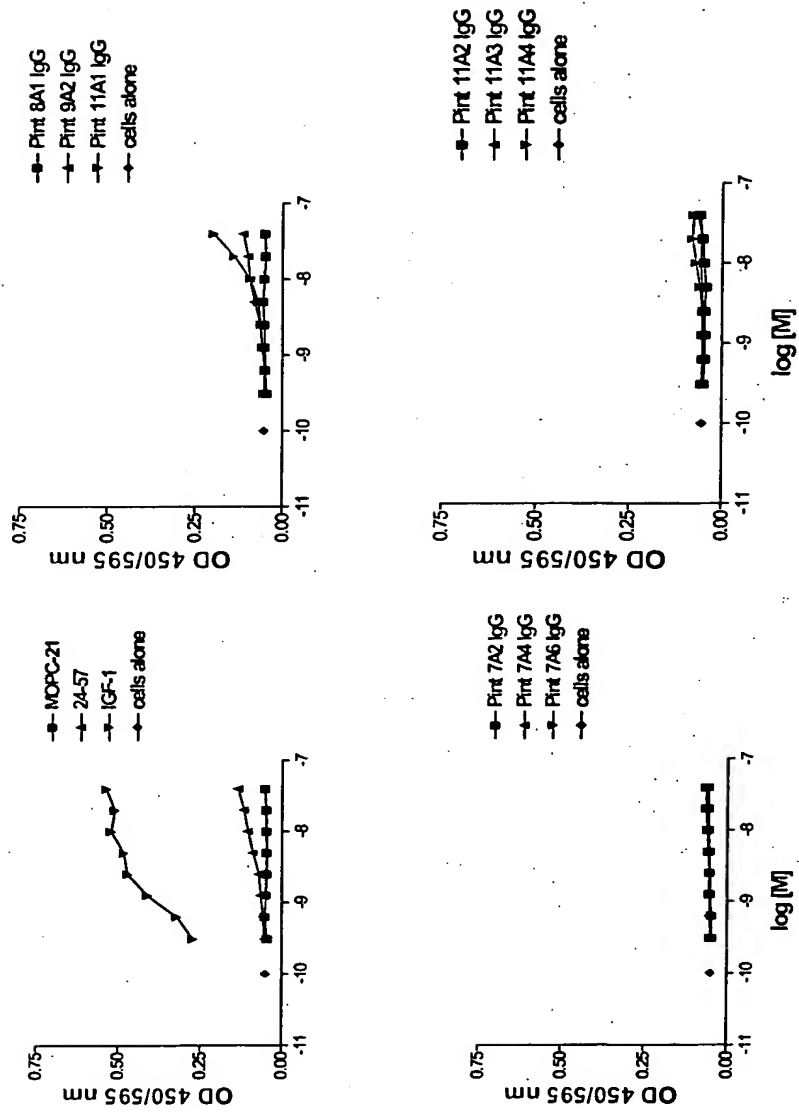


Figure 9

**IGF-1R mAbs – blockade of IGF-driven receptor
tyrosine phosphorylation – ELISA**

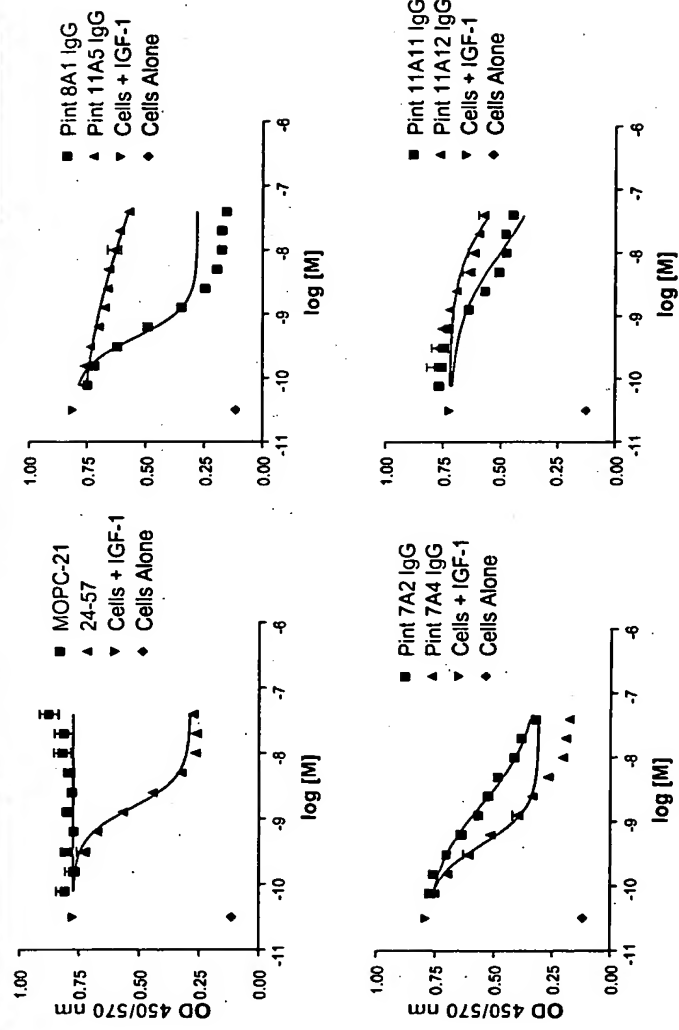


Figure 10

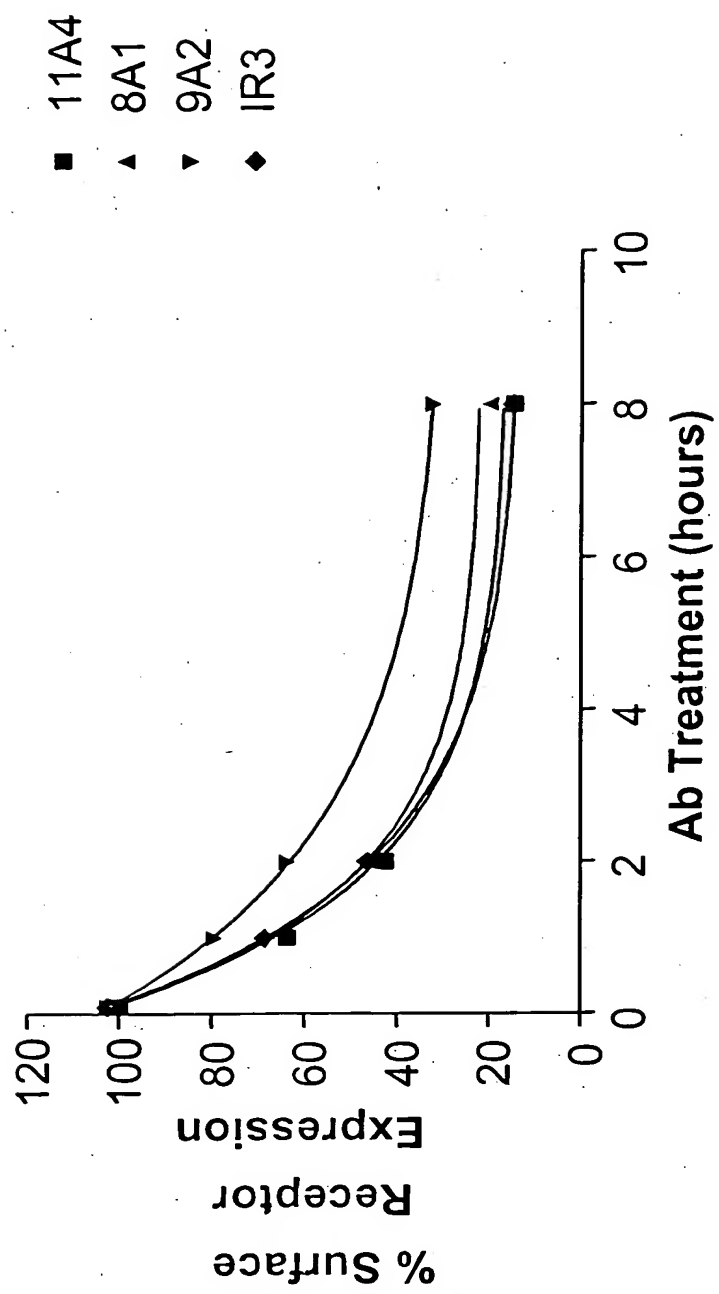


Figure 11

Time course of IGF-1R loss by IGF-1R mAbs,
but not w/ IGF-1 – NIH 3T3 hu-IGF-1R cells

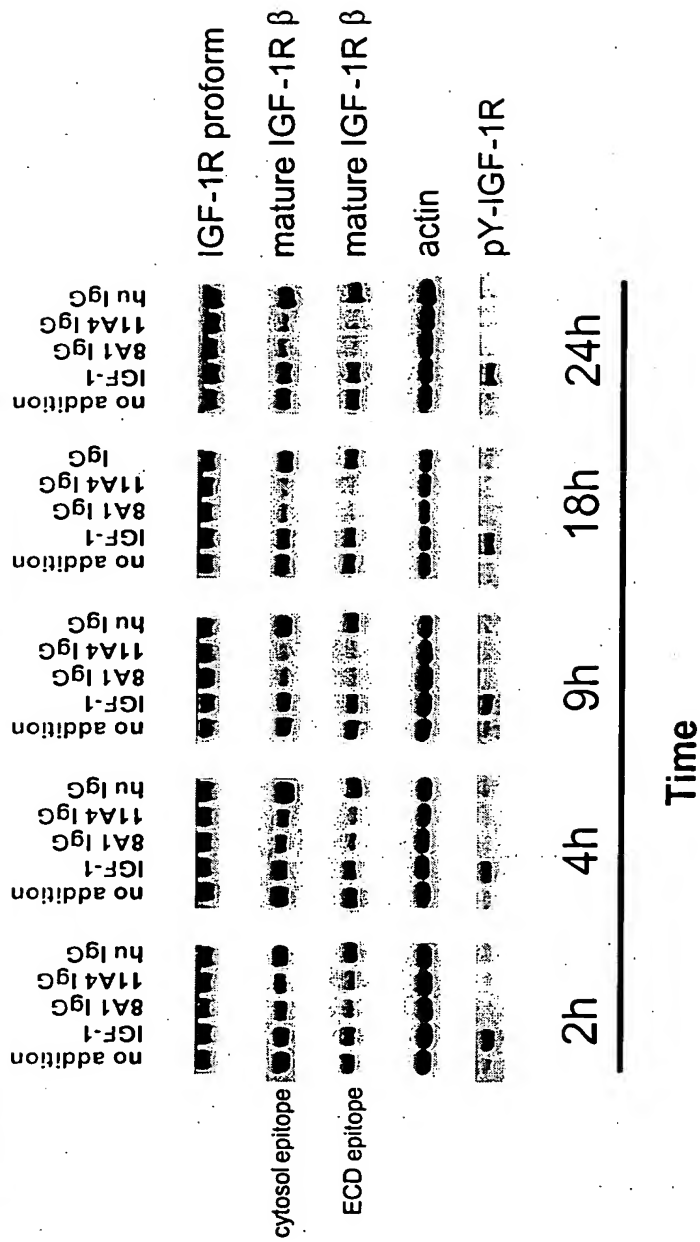


Figure 12

IGF-1R down-regulation *in vitro*
by IGF-1R mAbs – NIH 3T3 hu-IGF-1R cells

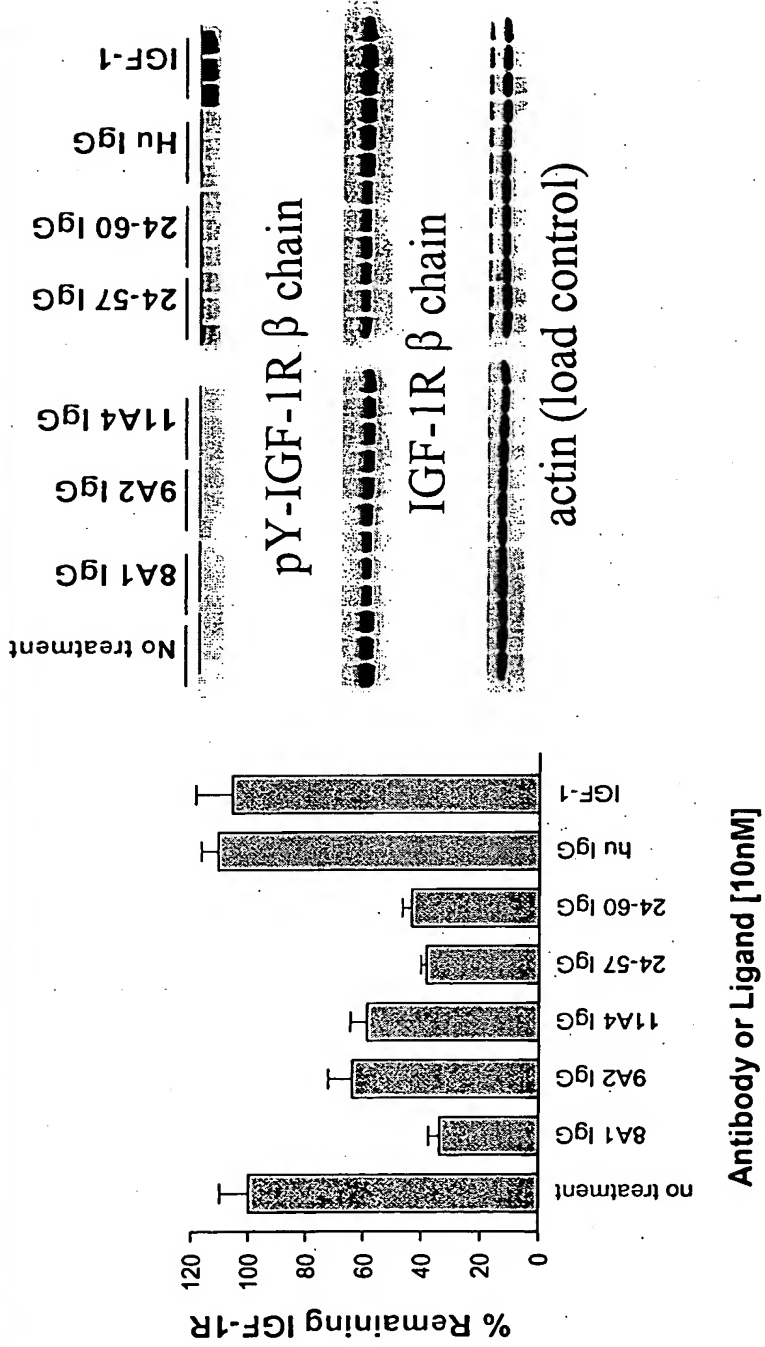


Figure 13

IGF-1R down-regulation *in vitro* by
IGF-1R mAbs – A549 NSCLC cells

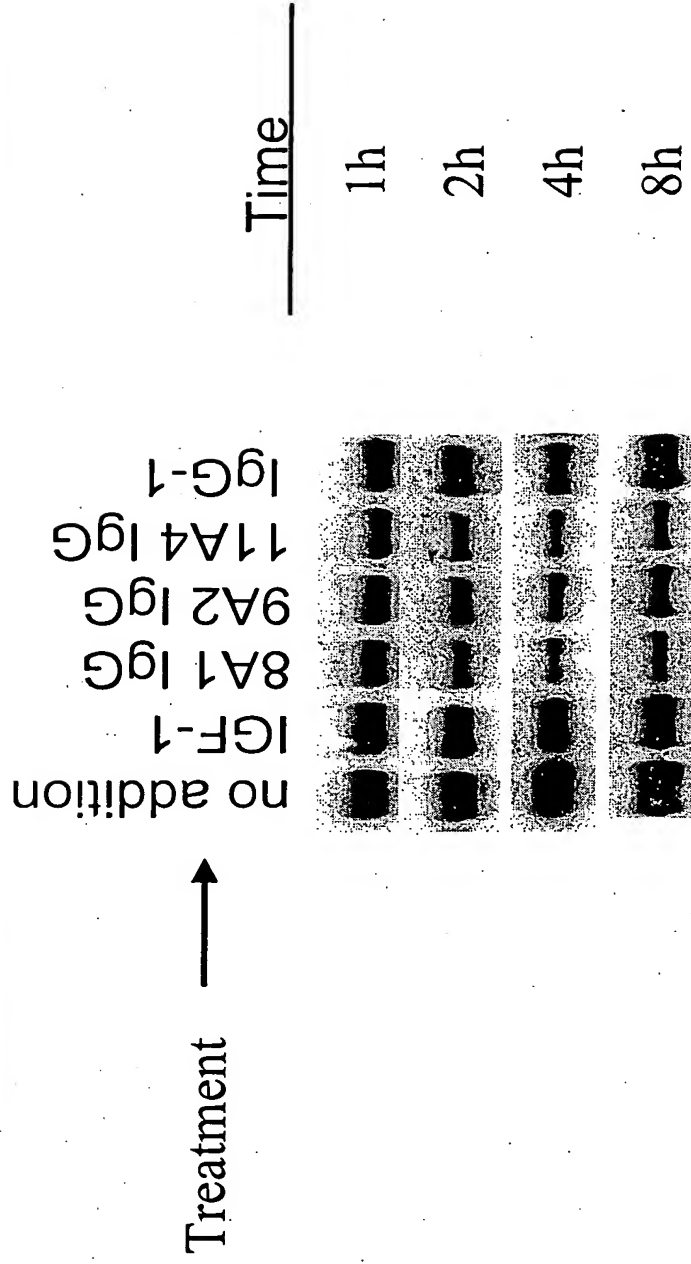
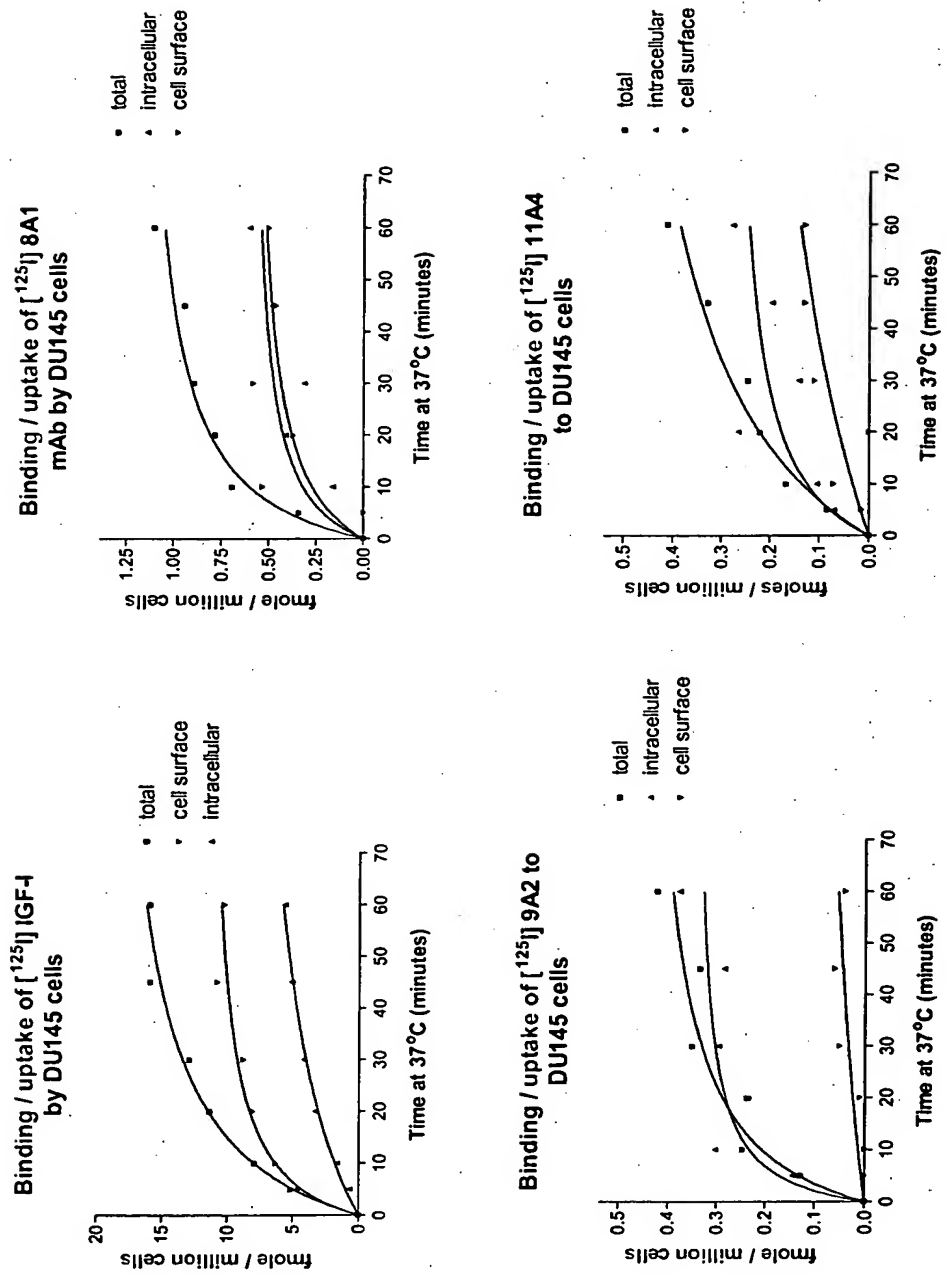


Figure 14



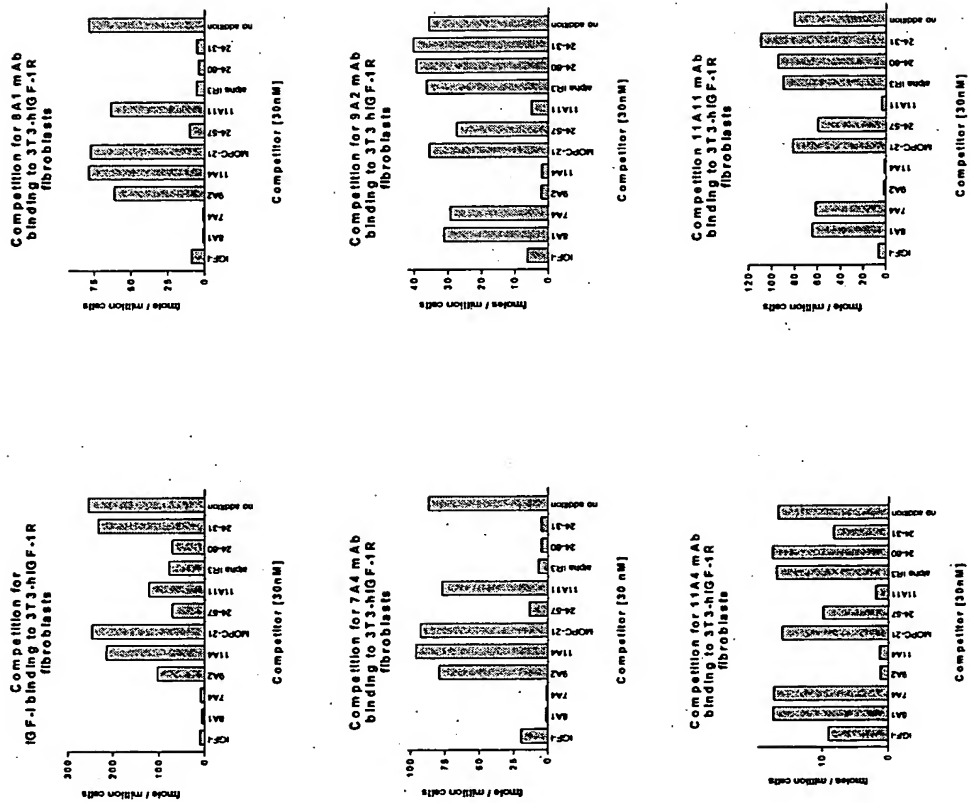


Figure 15

Figure 16

Distinct epitopes for mAbs

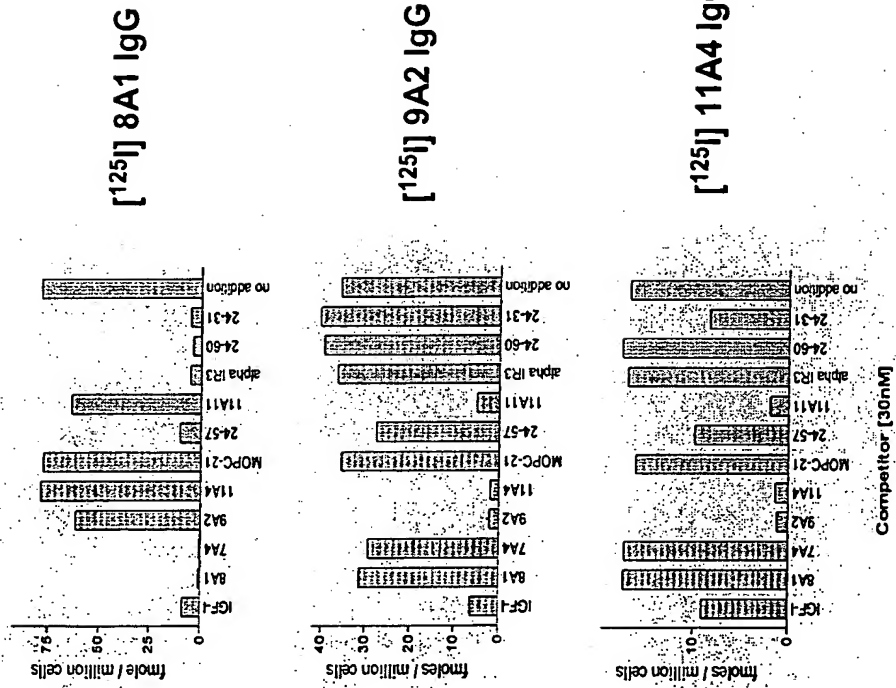


Figure 17

Inhibition of tumor growth/IGF-1R expression with
IGF-IR mAbs

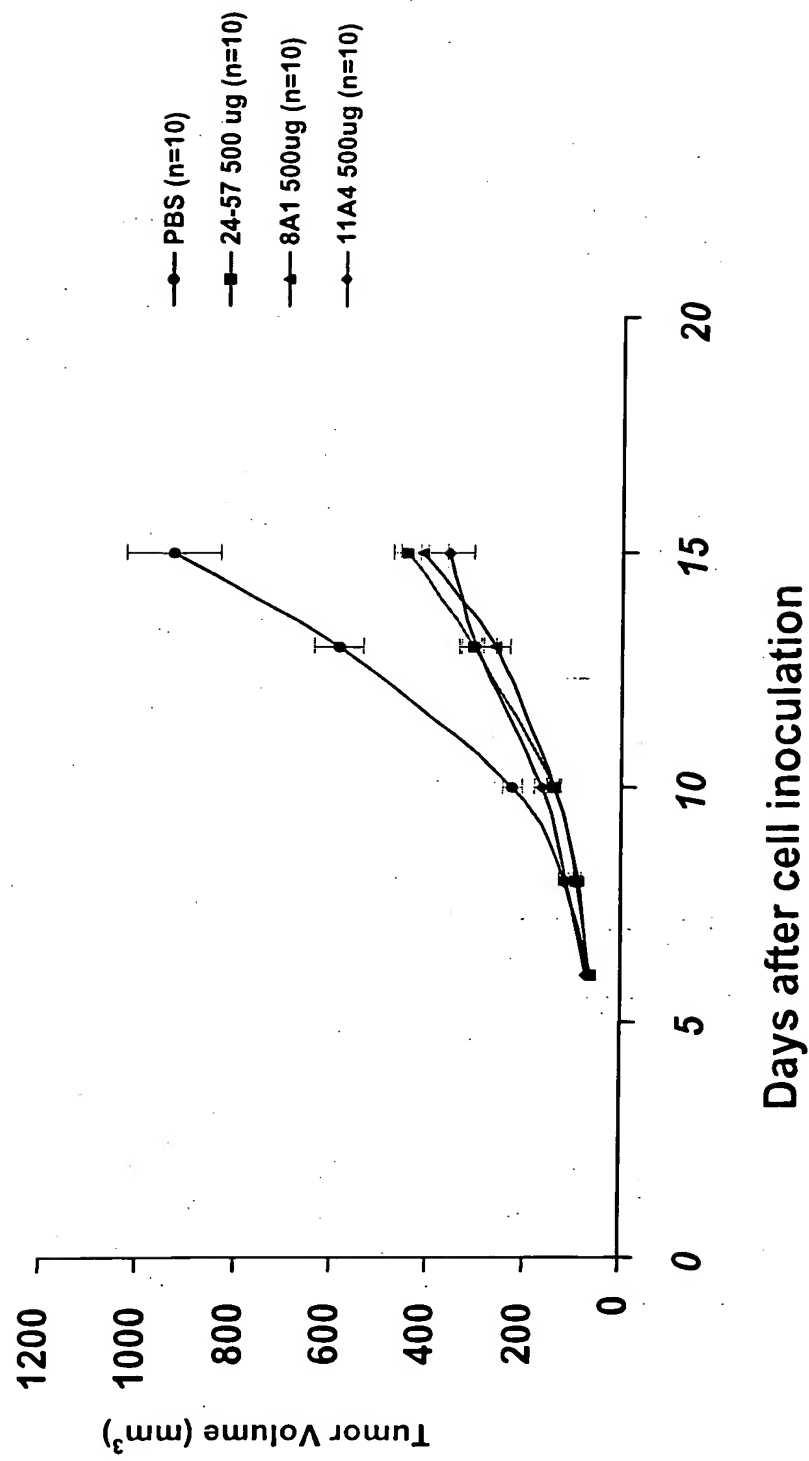


Figure 18

Inhibition of tumor growth/IgG-IR expression
with IGF-IR 8A1 IgG

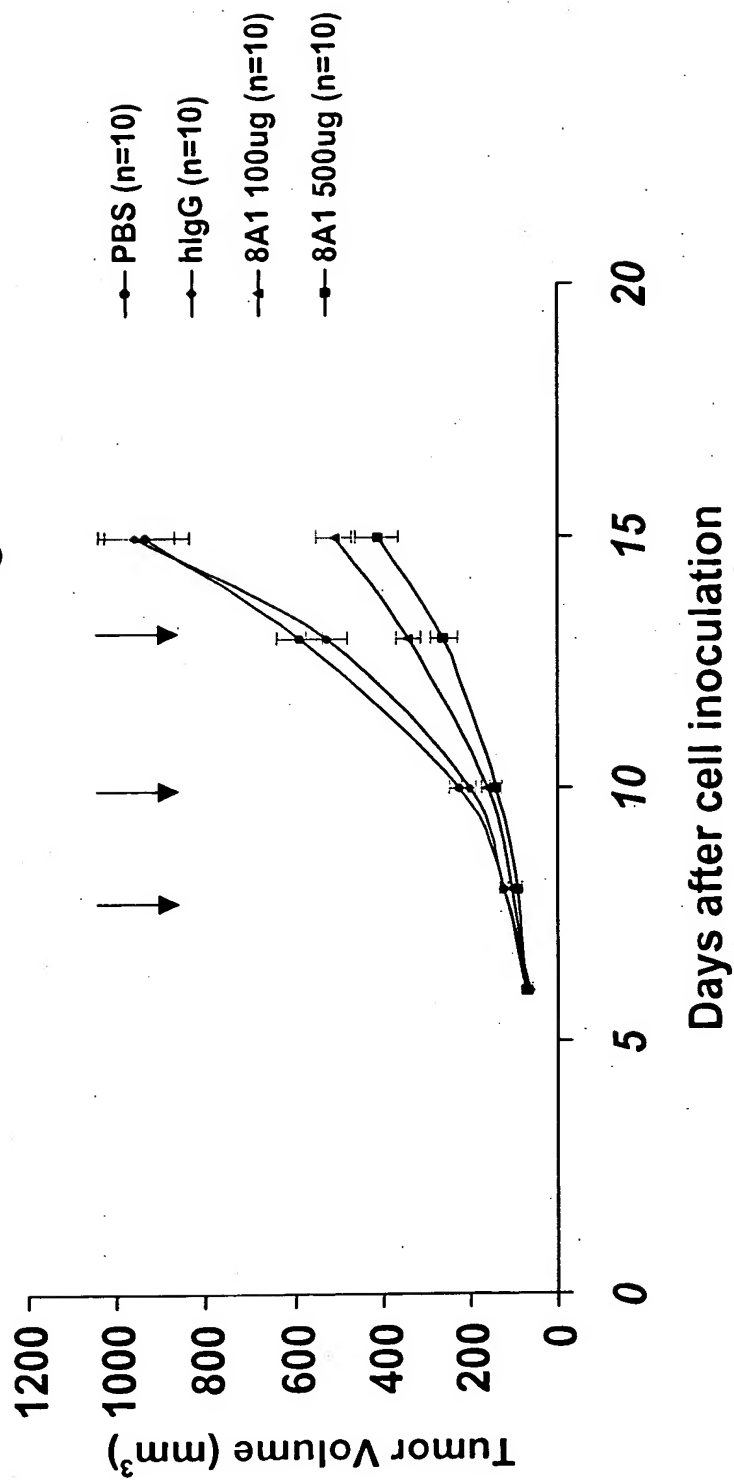


Figure 19

